

 \mathbf{B}

	LD50 (pfu)			
	WT	AV1	AV2	
IN		•		
Balb/c	1 × 10 ⁴	3 × 10 ⁸	2.5 × 10 ⁸	
CD-1	1 × 10 ⁸	2×10^{8}	nd	
IV				
CD-1	1 × 10 ⁸	8 × 10 ⁹	nd	

 \mathbf{C}

·	PFU	Morbidity	Mortality
WT VSV	10 ¹	3/3	3/3
AV2	10 ⁷	0/3	0/3
AV@(106) + WT VSV	10¹	0/3	0/3
	10 ²	0/3	0/3
·	10 ³	0/3	0/3

FIGURE 1A-C

WO 2004/085658

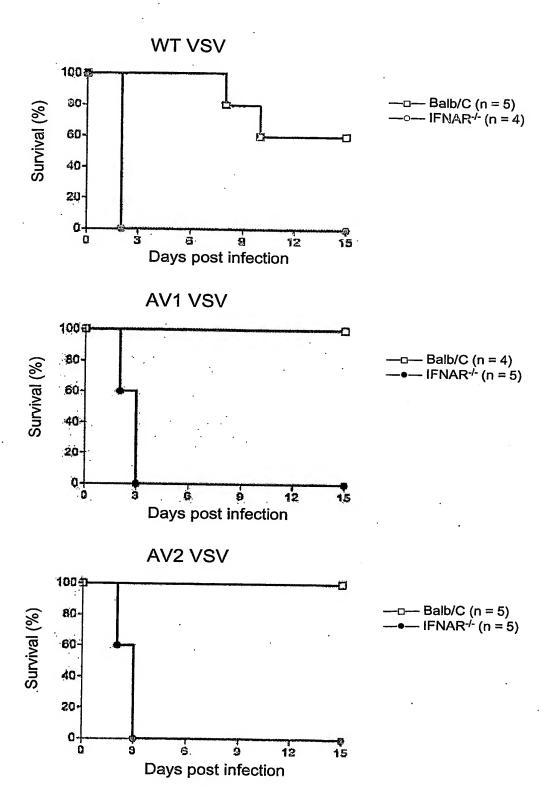


FIGURE 1D

WO 2004/085658

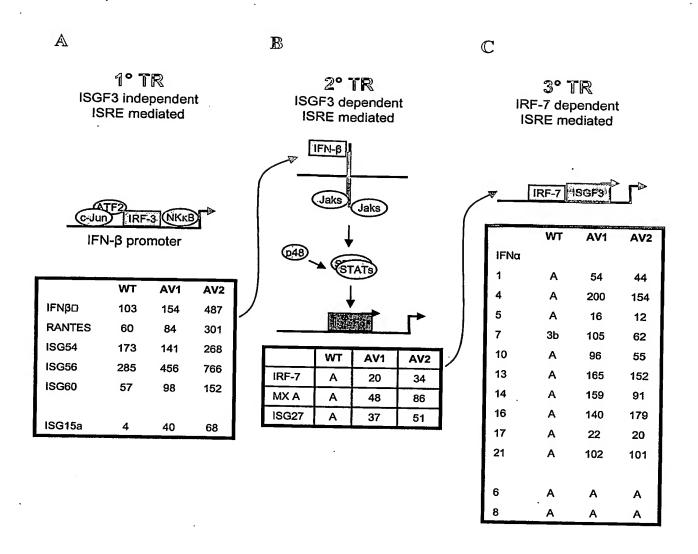
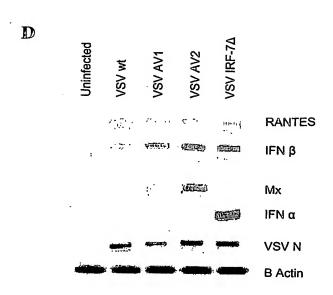


FIGURE 2A-C



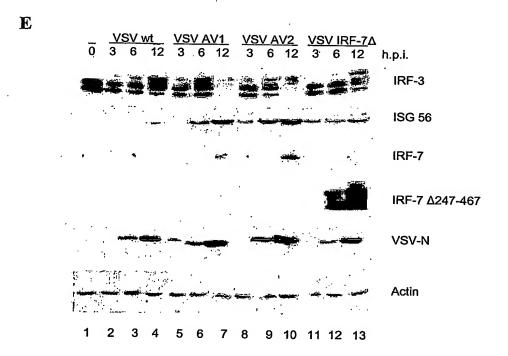


FIGURE 2D-E

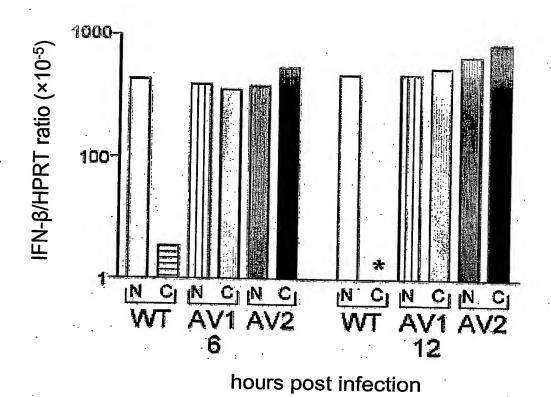
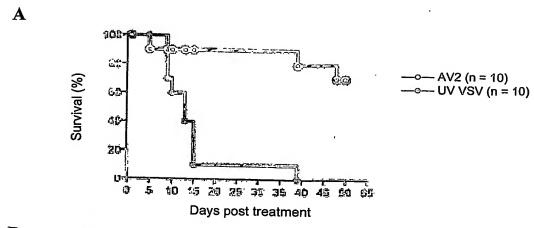
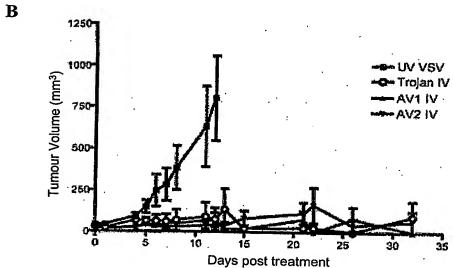


FIGURE 3





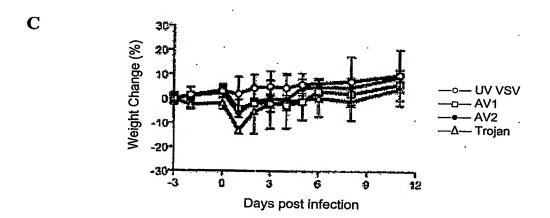
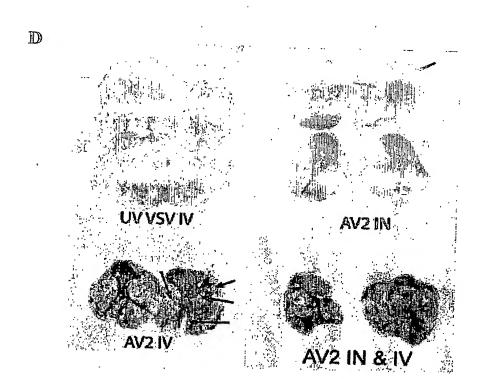


FIGURE 4A-C



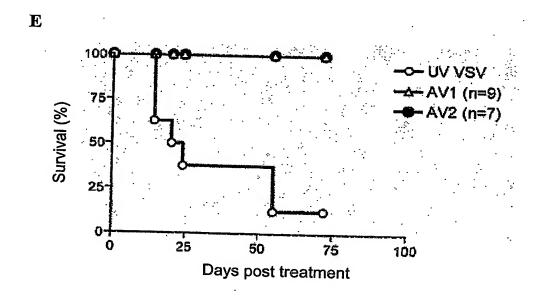
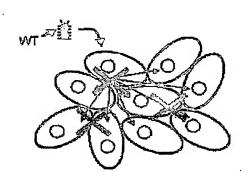


FIGURE 4D-E

Α



В

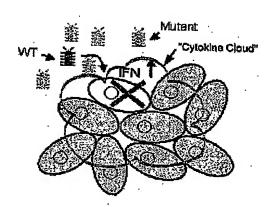


FIGURE 5

> EMDTHDPHQL AWVLDSVSHFKStop 50 51 52 53 54 55 56 57 58 59 221 226

Substitutions M51R

R

M51A

Α

M51-54A

AAAA

Deletions

M51

M51-54

M51-57

Mut2-like Mutants

M51R M51A M51-54A

ΔM51 ΔM51-54

ΔM51-57

Mut3-like Mutants

V221F S226R

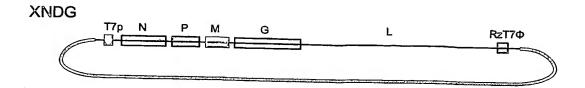
*S226R

*V221F

*ΔV221-S226

Compound Mutants

M51R & V221F S226R M51A & V221F S226R M51-54A & V221F S226R ΔM51 & V221F S226R ΔM51-54 & V221F S226R ΔM51-57 & V221F S226R



E M D T H D P H Q L 50 51 52 53 54 55 56 57 58 59

XNDG M4 E X D T H D P H Q L 50 X 52 53 54 55 56 57 58 59

E X X X X D P H Q L 550 X X X 555 56 57 58 59

FIGURE 7

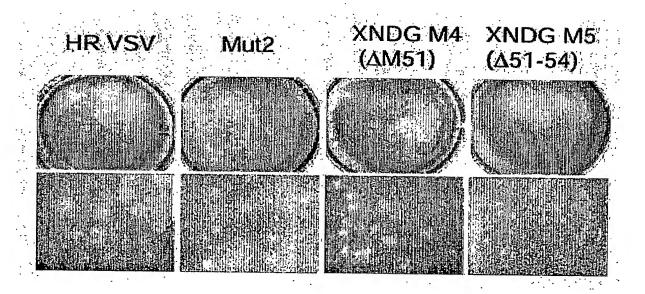


FIGURE 8

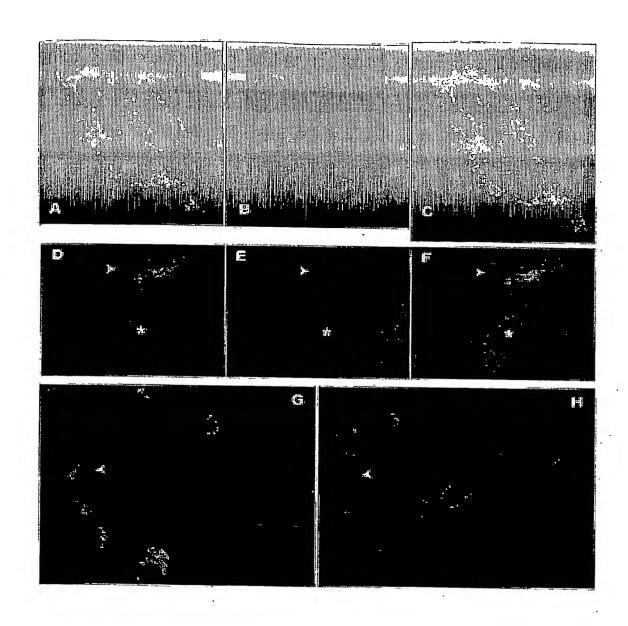
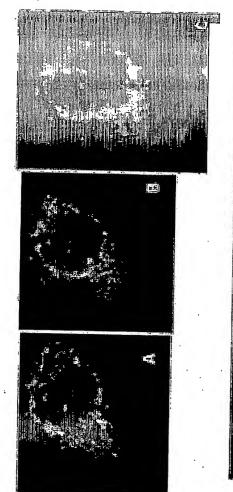
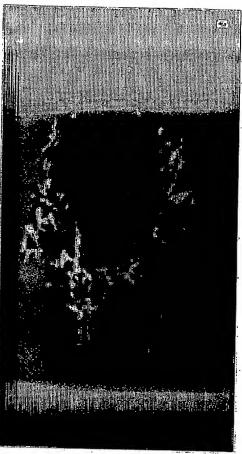


FIGURE 9A-H





Genome Sequence for VSV Mutant AV1

ACGAAGACAAACAAACCATTATTATCATTAAAAGGCTCAGGAGAAACTTTAACAGTAATCAAAATGTCTGTT ACAGTCAAGAGAATCATTGACAACACAGTCATAGTTCCAAAAACTTCCTGCAAATGAGGATCCAGTGGAATAC CCGGCAGATTACTTCAGAAAATCAAAGGAGATTCCTCTTTACATCAATACTACAAAAAGTTTGTCAGATCTA GCATTGAAGGACATCCGGGGTAAGTTGGATAAAGATTGGTCAAGTTTCGGAATAAACATCGGGAAGGCAGGG GATACAATCGGAATATTTGACCTTGTATCCTTGAAAGCCCTGGACGGTGTACTTCCAGATGGAGTATCGGAT GCTTCCAGAACCAGCGCAGATGACAAATGGTTGCCTTTGTATCTACTTGGCTTATACAGAGTGGGCAGAACA CAAATGCCTGAATACAGAAAAAGGCTCATGGATGGGCTGACAAATCAATGCAAAATGATCAATGAACAGTTT GAACCTCTTGTGCCAGAAGGTCGTGACATTTTTGATGTGTGGGGAAATGACAGTAATTACACAAAAATTGTC ${\tt GCTGCAGTGGACATGTTCCACATGTTCAAAAAACATGAATGTGCCTCGTTCAGATACGGAACTATTGTT}$ TCCAGATTCAAAGATTGTGCTGCATTGGCAACATTTGGACACCTCTGCAÁAATAACCGGAATGTCTACAGAA GATGTAACGACCTGGATCTTGAACCGAGAAGTTGCAGATGAGATGGTCCAAATGATGCTTCCAGGCCAAGAA $\tt GTCAAAAACCCTGCCTTCCACTTCTGGGGGCAATTGACAGCTCTTCTGCTCAGATCTACCAGAGCAAGGAAT$ ${\tt GCCCGACAGCCTGATGACATTGAGTATACATCTCTTACTACAGCAGGTTTGTTGTACGCTTATGCAGTAGGA}$ TCCTCTGCTGACTTGGCACAACAGTTTTGTGTTGGAGATAGCAAATACACTCCAGATGATAGTACCGGAGGA TTGACGACTAATGCACCGCCACAAGGCAGAGATGTGGTCGAATGGCTCGGATGGTTTGAAGATCAAAACAGA AAACCGACTCCTGATATGATGCAGTATGCGAAACGAGCAGTCATGTCACTGCAAGGCCTAAGAGAGAAGACA ATTGGCAAGTATGCTAAGTCAGAATTTGACAAATGACCCTATAATTCTCAGATCACCTATTATATATTATGC TACATATGAAAAAACTAACAGATATCATGGATAATCTCACAAAAGTTCGTGAGTATCTCAAGTCCTATTCT CGTCTAGATCAGGCGGTAGGAGAGATAGATGAGATCGAAGCACAACGAGCTGAAAAGTCCAATTATGAGTTG ${\tt TTCCAAGAGGACGGAGTGGAAGAGCATACTAGGCCCTCTTATTTTCAGGCAGCAGATGATTCTGACACAGAA}$ TCTGAACCAGAAATTGAAGACAATCAAGGCTTGTATGTACCAGATCCGGAAGCTGAGCAAGTTGAAGGCTTT ${\tt ATACAGGGGCCTTTAGATGACTATGCGGATGAGGACGTGGATGTTGTATTCACTTCGGACTGGAAACAGCCT}$ GAGCTTGAATCCGACGAGCATGGAAAGACCTTACGGTTGACATTGCCAGAGGGTTTAAGTGGAGAGCAGAAA ${ t TCCCAGTGGCTTTTGACGATTAAAGCAGTCGTTCAAAGTGCCAAACACTGGAATCTGGCAGAGTGCACATTT$ GAAGCATCGGGAGAAGGGGTCATCATAAAAAAGCGCCAGATAACTCCGGATGTATATAAGGTCACTCCAGTG CAACCCAAGAAAGCAAGTCTTCAGCCTCTCACCATATCCTTGGATGAATTGTTCTCATCTAGAGGAGAATTC ATCTCTGTCGGAGGTAACGGACGAATGTCTCATAAAGAGGCCATCCTGCTCGGTCTGAGGTACAAAAGTTG TACAATCAGGCGAGAGTCAAATATTCTCTGTAGACTATGAAAAAAAGTAACAGATATCACAATCTAAGTGTT ATCCCAATCCATTCATCATGAGTTCCTTAAAGAAGATTCTCGGTCTGAAGGGGAAAGGTAAGAAATCTAAGA AATTAGGGATCGCACCACCCCTTATGAAGAGGACACTAACATGGAGTATGCTCCGAGCGCTCCAATTGACA AATCCTATTTTGGAGTTGACGAGAGGGACACTCATGATCCGCATCAATTAAGATATGAGAAATTCTTCTTTA CAGTGAAAATGACGGTTAGATCTAATCGTCCGTTCAGAACATACTCAGATGTGGCAGCCGCTGTATCCCATT

 ${\tt GGGATCACATGTACATCGGAATGGCAGGGAAACGTCCCTTCTACAAGATCTTGGCTTTTTTGGGTTCTTCTA}$ ATCTAAAGGCCACTCCAGCGGTATTGGCAGATCAAGGTCAACCAGAGTATCACGCTCACTGTGAAGGCAGGG CTTATTTGCCACACAGAATGGGGAAGACCCCTCCCATGCTCAATGTACCAGAGCACTTCAGAAGACCATTCA ATATAGGTCTTTACAAGGGAACGGTTGAGCTCACAATGACCATCTACGATGATGAGCACTGGAAGCAGCTC CTATGATCTGGGATCATTTCAATTCTTCCAAATTTTCTGATTTCAGAGAGAAGGCCTTAATGTTTGGCCTGA TTGTCGAGAAAAAGGCATCTGGAGCTTGGGTCCTGGATTCTGTCAGCCACTTCAAATGAGCTAGTCTAGCTT CCAGCTTCTGAACAATCCCCGGTTTACTCAGTCTCTCCTAATTCCAGCCTTTCGAACAACTAATATCCTGTC ${\tt TTTTCTATCCCTATGAAAAAAACTAACAGAGATCGATCTGTTTCCTTGACACCATGAAGTGCCTTTTGTACT}$ ${\tt TAGCTTTTTATTCATCGGGGTGAATTGCAAGTTCACCATAGTTTTTCCATACAACCGAAAAGGAAACTGGA}$ AAAATGTTCCTTCCAATTACCATTATTGCCCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCA $\tt CAGCCTTACAAGTCAAAATGCCCAAGAGTCACAAGGCTATTCAAGCAGACGGTTGGATGTCATGCTTCCA$ ${\tt AATGGGTCACTACTTGTGATTTCCGCTGGTACGGACCGAAGTATATAACACATTCCATCCGATCCTTCACTC}$ $\tt CATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAACTTGGCTGAATCCAGGCTTCCCTC$ CTCAAAGTTGTGGATATGCAACTGTGACGGATGCTGAAGCAGCGATTGTCCAGGTGACTCCTCACCATGTGC TTGTTGATGAATACACAGGAGAATGGGTTGATTCACAGTTCATCAACGGAAAATGCAGCAATGACATATGCC CCACTGTCCATAACTCCACAACCTGGCATTCCGACTATAAGGTCAAAGGGCTATGTGATTCTAACCTCATTT CCATGGACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTAGGAAAGGAGGGCACAGGGTTCAGAA ${\tt TCCCATCAGGTGTCTGGTTCGAGATGGCTGATAAGGATCTCTTTGCTGCAGCCAGATTCCCTGAATGCCCAG}$ ${\tt AAGGGTCAAGTATCTCTGCTCCATCTCAGACCTCAGTGGATGTAAGTCTCATTCAGGACGTTGAGAGGATCT}$ ${\tt TGGATTATTCCCTCTGCCAAGAAACCTGGAGCAAAATCAGAGCGGGTCTTCCCATCTCTCCAGTGGATCTCA}$ GCTATCTTGCTCCTAAAAACCCAGGAACCGGTCCTGTCTTTACCATAATCAATGGTACCCTAAAATACTTTG AGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCTCAAGAATGGTCGGAATGATCAGTGGAACTA $\tt CCACAGAAAGGGGAACTGTGGGATGACTGGGGCTCCATATGAAGACGTGGAAATTGGACCCAATGGAGTTCTGA$ ${\tt GGACCAGTTCAGGATATAAGTTTCCTTTATATATGATTGGACATGGTATGTTGGACTCCGATCTTCATCTTA}$ ${\tt GCTCAAAGGCTCAGGTGTTTGAACATCCTCACATTCAAGACGCTGCTTCGCAGCTTCCTGATGATGAGACTT}$ GCTCTATTGCCTCTTTTTTCTTTATCATAGGGTTAATCATTGGACTATTCTTGGTTCTCCGAGTTGGTATTT ATCTTTGCATTAAATTAAAGCAĊACCAAGAAAAGACAGATTTATACAGACATAGAGATGAACCGACTTGGGA AGTAACTCAAATCCTGCACAACAGATTCTTCATGTTTGAACCAAATCAACTTGTGATATCATGCTCAAAGAG GCCTTAATTATATTTTAATTTTTAATTTTTATGAAAAAACTAACAGCAATCATGGAAGTCCACGATTTTGA GACCGACGAGTTCAATGATTCAATGAAGATGACTATGCCACAAGAGAATTCCTGAATCCCGATGAGCGCAT GACGTACTTGAATCATGCTGATTACAATTTGAATTCTCCTCTAATTAGTGATGATATTGACAATTTGATCAG ${\tt GAAATTCAATTCTCTTCCGATTCCCTCGATGTGGGATAGTAAGAACTGGGATGGAGTTCTTGAGATGTTAAC}$ ${\tt TCATGATGCCAGTCAAGGGTATAGTTTTTTACATGAAGTGGACAAAGAGGCAGAAATAACATTTGACGTGGT$

CAAAATTCTCGCTTATTTGTGTCAAAAGTTTTTGGACTTACACAAGTTGACATTAATCTTAAATGCTGTCTC ${ t ATGCAGGCTTAGGGTTCCCAGCTTGGGTCCTACTTTATTTCAGAAGGATGGGCTTACTTCAAGAAACTTGA}$ ${\tt TATTCTAATGGACCGAAACTTTCTGTTAATGGTCAAAGATGTGATTATAGGGAGGATGCAAACGGTGCTATC}$ CATGGTATGTAGAATAGACAACCTGTTCTCAGAGCAAGACATCTTCTCCCTTCTAAATATCTACAGAATTGG AGATAAAATTGTGGAGAGGCAGGGAAATTTTTCTTATGACTTGATTAAAATGGTGGAACCGATATGCAACTT GAGGCTGATGAAATTAGCAAGAGAATCAAGGCCTTTAGTCCCACAATTCCCTCATTTTGAAAATCATATCAA GACTTCTGTTGATGAAGGGGCAAAAATTGACCGAGGTATAAGATTCCTCCATGATCAGATAATGAGTGTGAA ${\tt AACAGTGGATCTCACACTGGTGATTTATGGATCGTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTTATACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGCTTTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGATTATAGAT$ ${\tt AAGTGATTTAGCTCGGATTGTTCTATTTCAACAGTTCAATGATCATAAAAAGTGGTTCGTGAATGGAGACTT}$ GCTCCCTCATGATCATCCCTTTAAAAGTCATGTTAAAGAAAATACATGGCCCACAGCTGCTCAAGTTCAAGA ${\tt TTTTGGAGATAAATGGCATGAACTTCCGCTGATTAAATGTTTTGAAATACCCGACTTACTAGACCCATCGAT}$ TCCTATCCCTAGTAAAAAGGTGTTGCAGACTATGTTGGACACAAAGGCTACCAATTGGAAAGAATTTCTTAA ${\tt GTTGGCAGGTAGATTTTTCTCCCTAATGTCTTGGAAATTGCGAGAATACTTTGTAATTACCGAATATTTGAT}$ AAAGACTCATTTCGTCCCTATGTTTAAAGGCCTGACAATGGCGGACGATCTAACTGCAGTCATTAAAAAGAT GTŁAGATTCCTCATCCGGCCAAGGATTGAAGTCATATGAGGCAATTTGCATAGCCAATCACATTGATTACGA AAAATGGAATAACCACCAAAGGAAGTTATCAAACGGCCCAGTGTTCCGAGTTATGGGCCAGTTCTTAGGTTA TCCATCCTTAATCGAGAGACTCATGAATTTTTTGAGAAAAGTCTTATATACTACAATGGAAGACCAGACTT GATGCGTGTTCACAACAACACACTGATCAATTCAACCTCCCAACGAGTTTGTTGGCAAGGACAAGAGGGTGG ACTGGAAGGTCTACGGCAAAAAGGATGGAGTATCCTCAATCTACTGGTTATTCAAAGAGAGGGCTAAAATCAG AAACACTGCTGTCAAAGTCTTGGCACAAGGTGATAATCAAGTTATTTGCACACAGTATAAAACGAAGAAATC GAGAAACGTTGTAGAATTACAGGGTGCTCTCAATCAAATGGTTTCTAATAATGAGAAAATTATGACTGCAAT ${ t CAAAATAGGGACAGGGAAGTTAGGACTTTTGATAAATGACGATGAGACTATGCAATCTGCAGATTACTTGAA}$ TTATGGAAAAATACCGATTTTCCGTGGAGTGATTAGAGGGTTAGAGACCAAGAGATGGTCACGAGTGACTTG TGTCACCAATGACCAAATACCCACTTGTGCTAATATAATGAGCTCAGTTTCCACAAATGCTCTCACCGTAGC TCATTTTGCTGAGAACCCAATCAATGCCATGATACAGTACAATTATTTTGGGACATTTGCTAGACTCTTGTT GATGATGCATGATCCTGCTCTTCGTCAATCATTGTATGAAGTTCAAGATAAGATACCGGGCTTGCACAGTTC TTTGATTAGAGCCTTCCCAGATCCCGTAACAGAAAGTCTCTCATTCTGGAGATTCATCCATGTaCATGCTCG AAGTGAGCATCTGAAGGAGATGAGTGCAGTATTTGGAAACCCCGAGATAGCCAAGTTTCGAATAACTCACAT AGACAAGCTAGTAGAAGATCCAACCTCTCTGAACATCGCTATGGGAATGAGTCCAGCGAACTTGTTAAAGAC TGAGGTTAAAAAATGCTTAATCGAATCAAGACAAACCATCAGGAACCAGGTGATTAAGGATGCAACCATATA

TTTGTATCATGAAGAGGATCGGCTCAGAAGTTTCTTATGGTCAATAAATCCTCTGTTCCCTAGATTTTTAAG TCGGAACTCCTTTAAGAAAAAGTATCATAGGGAATTGGATGATTTGATTGTGAGGAGTGAGGTATCCTCTTT GACACATTTAGGGAAACTTCATTTGAGAAGGGGATCATGTAAAATGTGGACATGTTCAGCTACTCATGCTGA ${\tt TCCACAACATCGAAAAGAGACTCCTTGTGCACCATGTAACACATCAGGGTTCAATTATGTTTCTGTGCATTG$ ${\tt TCCAGACGGGATCCATGACGTCTTTAGTTCACGGGGACCATTGCCTGCTTATCTAGGGTCTAAAACATCTGA}$ ATCTACATCTATTTTGCAGCCTTGGGAAAGGAAAGCAAAGTCCCACTGATTAAAAGAGCTACACGTCTTAG ${\tt AGATGCTATCTCTTGGTTGAACCCGACTCTAAACTAGCAATGACTATACTTTCTAACATCCACTCTTT}$ ${\tt ATCTCGGATGAGCCATGGTGGGTTCGCATCTCAGAGCACTGCAGCATTGACCAGGTTGATGGCAACTACAGA}$ $\tt CACCATGAGGGATCTGGGAGATCAGAATTTCGACTTTTTATTCCAGGCAACGTTGCTCTATGCTCAGATTAC$ GAGACCCATAGAAGAGATCACCCTGGACTCAAGTATGGACTACACGCCCCCAGATGTATCCCATGTGCTGAA ${\tt GACATGGAGGAATGGGGAAGGGTTCGTGGGGACAAGAGATAAAACAGATCTATCCTTTAGAAGGGAATTGGAA}$ GAATTTAGCACCTGCTGAGCAATCCTATCAAGTCGGCAGATGTATAGGTTTTCTATATGGAGACTTGGCGTA TTTCTTAAAAGGGTTGCTAGACGGATTAATGAGAGCAAGTTGCTGCCAAGTAATACACCGGAGAAGTCTGGC $\tt CTCCTATCCGACAAGCAACCGTGATATGGGGGGTGATTGTCAGAAATACCTTCAAATACCAATGCCGTCTAAT$ TGAAAAGGGAAAATACAGATCACATTATTCACAATTATGGTTATTCTCAGATGTCTTATCCATAGACTTCAT TGGACCATTCTCTATTTCCACCACCCTCTTGCAAATCCTATACAAGCCATTTTTATCTGGGAAAGATAAGAA ATTCTTCACCAAGGACATATTATTGTGTCCAGAGGAAATCAGACATGCTTGCAAGTTCGGGATTGCTAAGGA TAATAATAAAGACATGAGCTATCCCCCTTGGGGAAGGGAATCCAGAGGGACAATTACAACAATCCCTGTTTA ${\tt CAGGTTGGGCCAGTTACCAACTGGCGCTCATTATAAAATTCGGAGTATATTACATGGAATGGGAATCCATTA}$ TGCCCTAGAAACTTTAGGAGGAGATAAATCGAGATGTGTAAATGGTGAAACATGTTGGGAATATCCATCTGA $\tt CTTATGTGACCCAAGGACTTGGGACTATTTCCTCCGACTCAAAGCAGGCTTGGGGCTTCAAATTGATTTAAT$ TGTAATGGATATGGAAGTTCGGGGATTCTTCTACTAGCCTGAAAATTGAGACGAATGTTAGAAATTATGTGCA ${\tt CCGGATTTTGGATGAGCAAGGAGTTTTAATCTACAAGACTTATGGAACATATATTTGTGAGAGCGAAAAGAA}$ ${\tt TGCAGTAACAATCCTTGGTCCCATGTTCAAGACGGTCGACTTAGTTCAAACAGAATTTAGTAGTTCTCAAAC}$ GTCTGAAGTATATATGGTATGTAAAGGTTTGAAGAAATTAATCGATGAACCCAATCCCGATTGGTCTTCCAT

Nucleic Acid Sequence of the M Protein Gene for VSV Mutant AV1

Amino Acid Sequence for the M Protein of VSV Mutant AV1

 ${\tt MSSLKKILGLKGKGKKSKKLGIAPPPYEEDTNMEYAPSAPIDKSYFGVDERDTHDPHQLRYEKFFFTVKMTV} RSNRPFRTYSDVAAAVSHWDHMYIGMAGKRPFYKILAFLGSSNLKATPAVLADQGQPEYHAHCEGRAYLPHR MGKTPPMLNVPEHFRRPFNIGLYKGTVELTMTIYDDESLEAAPMIWDHFNSSKFSDFREKALMFGLIVEKKA SGAWVLDSVSHFK.$

Genome sequence for VSV Mutant AV2

ACGAAGACAAACAAACCATTATTATCATTAAAAGGCTCAGGAGAAACTTTAACAGTAATCAAAATGTCTGTT ACAGTCAAGAGAATCATTGACAACACAGTCATAGTTCCAAAAACTTCCTGCAAATGAGGATCCAGTGGAATAC CCGGCAGATTACTTCAGAAAATCAAAGGAGATTCCTCTTTACATCAATACTACAAAAAGTTTGTCAGATCTA GCATTGAAGGACATCCGGGGTAAGTTGGATAAAGATTGGTCAAGTTTCGGAATAAACATCGGGAAGGCAGGG GATACAATCGGAATATTTGACCTTGTATCCTTGAAAGCCCTGGACGGTGTACTTCCAGATGGAGTATCGGAT GCTTCCAGAACCAGCGCAGATGACAAATGGTTGCCTTTGTATCTACTTGGCTTATACAGAGTGGGCAGAACA CAAATGCCTGAATACAGAAAAAGGCTCATGGATGGGCTGACAAATCAATGCAAAATGATCAATGAACAGTTT GAACCTCTTGTGCCAGAAGGTCGTGACATTTTTGATGTGTGGGGAAATGACAGTAATTACACAAAAATTGTC GCTGCAGTGGACATGTTCTTCCACATGTTCAAAAAACATGAATGTGCCTCGTTCAGATACGGAACTATTGTT TCCAGATTCAAAGATTGTGCTGCATTGGCAACATTTGGACACCTCTGCAAAATAACCGGAATGTCTACAGAA GATGTAACGACCTGGATCTTGAACCGAGAAGTTGCAGATGAGATGGTCCAAATGATGCTTCCAGGCCAAGAA ATTGACAAGGCCGATTCATACATGCCTTATTTGATCGACTTTGGATTGTCTTCTAAGTCTCCATATTCTTCC GTCAAAAACCCTGCCTTCCACTTCTGGGGGCAATTGACAGCTCTTCTGCTCAGATCCACCAGAGCAAGGAAT ${\tt GCCCGACAGCCTGATGACATTGAGTATACATCTCTTACTACAGCAGGTTTGTTGTACGCTTATGCAGTAGGA}$ ${\tt TCCTCTGCTGACTTGGCACAACAGTTTTGTGTTGGAGATAGCAAATACACTCCAGATGATAGTACCGGAGGA}$ ${\tt TTGACGACTAATGCACCGCCACAAGGCAGAGATGTGGTCGAATGGCTCGGATGGTTTGAAGATCAAAACAGA}$ AAACCGACTCCTGATATGATGCAGTATGCGAAACGAGCAGTCATGTCACTGCAAGGCCTAAGAGAGAAGACA ATTGGCAAGTATGCTAAGTCAGAATTTGACAAATGACCCTATAATTCTCAGATCACCTATTATATATTATGC TACATATGAAAAAACTAACAGATATCATGGATAATCTCACAAAAGTTCGTGAGTATCTCAAGTCCTATTCT CGTCTAGATCAGGCGGTAGGAGATAGATGAGATCGAAGCACAACGAGCTGAAAAGTCCAATTATGAGTTG TTCCAAGAGGACGGAGTGGAAGAGCATACTAGGCCCTCTTATTTTCAGGCAGCAGATGATTCTGACACAGAA TCTGAACCAGAAATTGAAGACAATCAAGGCTTGTATGTACCAGATCCGGAAGCTGAGCAAGTTGAAGGCTTT ${\tt ATACAGGGGCCTTTAGATGACTATGCGGATGAGGACGTGGATGTTGTATTCACTTCGGACTGGAAACAGCCT}$ GAGCTTGAATCCGACGAGCATGGAAAGACCTTACGGTTGACATTGCCAGAGGGTTTAAGTGGAGAGCAGAAA ${\tt TCCCAGTGGCTTTTGACGATTAAAGCAGTCGTTCAAAGTGCCAAACACTGGAATCTGGCAGAGTGCACATTT}$ GAAGCATCGGGAGAAGGGGTCATCATAAAAAAGCGCCAGATAACTCCGGATGTATATAAGGTCACTCCAGTG ATGAACACACCGTCCCAATCGGAAGCCGTATCAGATGTTTGGTCTCTCAAAGACATCCATGACTTTC CAACCCAAGAAAGCAAGTCTTCAGCCTCTCACCATATCCTTGGATGAATTGTTCTCATCTAGAGGAGAATTC ATCTCTGTCGGAGGTAACGGACGAATGTCTCATAAAGAGGCCATCCTGCTCGGTCTGAGGTACAAAAAGTTG TACAATCAGGCGAGAGTCAAATATTCTCTGTAGACTATGAAAAAAAGTAACAGATATCACAATCTAAGTGTT ATCCCAATCCATTCATCATGAGTTCCTTAAAGAAGATTCTCGGTCTGAAGGGGAAAGGTAAGAAATCTAAGA AATTAGGGATCGCACCACCCCTTATGAAGAGGACACTAACATGGAGTATGCTCCGAGCGCTCCAATTGACA AATCCTATTTTGGAGTTGACGAGATGGACACTCATGATCCGCATCAATTAAGATATGAGAAATTCTTCTTTA CAGTGAAAATGACGGTTAGATCTAATCGTCCGTTCAGAACATACTCAGATGTGGCAGCCGCTGTATCCCATT

Figure 14

ATCTAAAGGCCACTCCAGCGGTATTGGCAGATCAAGGTCAACCAGAGTATCACGCTCACTGTGAAGGCAGGG $\tt CTTATTTGCCACACAGAATGGGGAAGACCCCTCCCATGCTCAATGTACCAGAGCACTTCAGAAGACCATTCA$ ATATAGGTCTTTACAAGGGAACGGTTGAGCTCACAATGACCATCTACGATGATGAGTCACTGGAAGCAGCTC TTGTCGAGAAAAAGGCATCTGGAGCTTGGTTCCTGGATTCTGTCAGACACTTCAAATGAGCTAGTCTAGCTT CCAGCTTCTGAACAATCCCCGGTTTACTCAGTCTCTCCTAATTCCAGCCTTTCGAACAACTAATATCCTGTC TTTTCTATCCCTATGAAAAAACTAACAGAGATCGATCTGTTTCCTTGACACCATGAAGTGCCTTTTGTACT TAGCTTTTTTATTCATCGGGGTGAATTGCAAGTTCACCATAGTTTTTCCATACAACCAAAAAGGAAACTGGA AAAATGTTCCTTCCAATTACCATTATTGCCCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCA CAGCCTTACAAGTCAAAATGCCCAAGAGTCACAAGGCTATTCAAGCAGACGGTTGGATGTGATGCTTCCA AATGGGTCACTACTTGTGATTTCCGCTGGTACGGACCGAAGTATATAACACATTCCATCCGATCCTTCACTC ${\tt CATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAACTTGGCTGAATCCAGGCTTCCCTC}$ CTCAAAGTTGTGGATATGCAACTGTGACGGATGCTGAAGCAGCGATTGTCCAGGTGACTCCTCACCATGTGC $\tt TTGTTGATGAATACACAGGAGAATGGGTTGATTCACAGTTCATCAACGGAAAATGCAGCAATGACATATGCC$ $\tt CCACTGTCCATAACTCCACAACCTGGCATTCCGACTATAAGGTCAAAGGGCTATGTGATTCTAACCTCATTT$ CCATGGACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTAGGAAAGGAGGGCACAGGGTTCAGAA GTAACTACTTTGCTTATGAAACTGGAGACAAGGCCTGCAAAATGCAGTACTGCAAGCGTTGGGGAGTCAGAC TCCCATCAGGTGTATGGTTCGAGATGGCTGATAAGGATCTCTTTGCTGCAGCCAGATTCCCTGAATGCCCAG AAGGGTCAAGTATCTCTGCTCCATCTCAGACCTCAGTGGATGTAAGTCTCATTCAGGACGTTGAGAGGATCT TGGATTATTCCCTCTGCCAAGAAACCTGGAGCAAAATCAGAGCGGGTCTTCCCATCTCTCCAGTGGATCTCA GCTATCTTGCTCCTAAAAACCCCAGGAACCGGTCCTGTCTTTACCATAATCAATGGTACCCTAAAATACTTTG AGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCTCAAGAATGGTCGGAATGATCAGTGGAACTA $\tt CCACAGAAAGGGAACTGTGGGATGACTGGGCTCCATATGAAGACGTGGAAATTGGACCCAATGGAGTTCTGA$ ${\tt GGACCAGTTCAGGATATAAGTTTCCTTTATATATGATTGGACATGGTATGTTGGACTCCGATCTTCATCTTA}$ GCTCAAAGGCTCAGGTGTTTGAACATCCTCACATTCAAGACGCTGCTGCGCAGCTTCCTGATGATGAGACTT GCTCTATTGCCTCTTTTTTCTTTATCATAGGGTTAATCATTGGACTATTCTTGGTTCTCCGAGTTGGTATTT ATCTTTGCATTAAATTAAAGCACACCAAGAAAAGACAGATTTATACAGACATAGAGATGAACCGACTTGGGA AGTAACTCAAATCCTGCACAACAGATTCTTCATGTTTGAACCAAATCAACTTGTGATATCATGCTCAAAGAG GCCTTAATTATATTTTAATTTTTAATTTTTATGAAAAAACTAACAGCAATCATGGAAGTCCACGATTTTGA ${\tt GACCGACGAGTTCAATGATTCAATGAAGATGACTATGCCACAAGAGAATTCCTGAATCCCGATGAGCGCAT}$ GACGTACTTGAATCATGCTGATTACAATTTGAATTCTCCTCTAATTAGTGATGATATTGACAATTTGATCAG GAAATTCAATTCTCTTCCGATTCCCTCGATGTGGGATAGTAAGAACTGGGATGGAGTTCTTGAGATGTTAAC TCATGATGCCAGTCAAGGGTATAGTTTTTTACATGAAGTGGACAAAGAGGCAGAAATAACATTTGACGTGGT

Figure 14 continued

CAAAATTCTCGCTTATTTGTGTCAAAAGTTTTTGGACTTACACAAGTTGACATTAATCTTAAATGCTGTCTC ${\tt ATGCAGGCTTAGGGTTCCCAGCTTGGGTCCTACTTTTATTTCAGAAGGATGGGCTTACTTCAAGAAACTTGA}$ TATTCTAATGGACCGAAACTTTCTGTTAATGGTCAAAGATGTGATTATAGGGAGGATGCAAACGGTGCTATC CATGGTATGTAGAATAGACAACCTGTTCTCAGAGCAAGACATCTTCTCCCTTCTAAATATCTACAGAATTGG AGATAAAATTGTGGAGAGGCAGGAAATTTTTCTTATGACTTGATTAAAATGGTGGAACCGATATGCAACTT GAAGCTGATGAAATTAGCAAGAGAATCAAGGCCTTTAGTCCCACAATTCCCTCATTTTGAAAATCATATCAA GACTTCTGTTGATGAAGGGGCAAAAATTGACCGAGGTATAAGATTCCTCCATGATCAGATAATGAGTGTGAA AACAGTGGATCTCACACTGGTGATTTATGGATCGTTCAGACATTGGGGTCATCCTTTTATAGATTATTACGC ${ t AAGTGATTTAGCTCGGATTGTTCTATTTCAACAGTTCAATGATCATAAAAAGTGGTTCGTGAATGGAGACTT}$ GCTCCCTCATGATCATCCCTTTAAAAGTCATGTTAAAGAAAATACATGGCCCACAGCTGCTCAAGTTCAAGA TTTTGGAGATAAATGGCATGAACTTCCGCTGATTAAATGTTTTGAAATACCCGACTTACTAGACCCATCGAT TCCTATCCCTAGTAAAAAGGTGTTGCAGACTATGTTGGACACAAAGGCTACCAATTGGAAAGAATTTCTTAA AGAGATTGATGAGAAGGGCTTAGATGATGATCTAATTATTGGTCTTAAAGGAAAGGAGAGGGAACTGAA GTTGGCAGGTAGATTTTTCTCCCTAATGTCTTGGAAATTGCGAGAATACTTTGTAATTACCGAATATTTGAT AAAGACTCATTTCGTCCCTATGTTTAAAGGCCTGACAATGGCGGACGATCTAACTGCAGTCATTAAAAAGAT GTTAGATTCCTCATCCGGCCAAGGATTGAAGTCATATGAGGCAATTTGCATAGCCAATCACATTGATTACGA ${\tt AAAATGGAATAACCACCAAAGGAAGTTATCAAACGGCCCAGTGTTCCGAGTTATGGGCCAGTTCTTAGGTTA}$ ${\tt TCCATCCTTAATCGAGAGACTCATGAATTTTTTTGAGAAAAGTCTTATATACTACAATGGAAGACCAGACTT}$ GATGCGTGTTCACAACAACACACTGATCAATTCAACCTCCCAACGAGTTTGTTGGCAAGGACAAGAGGGTGG ACTGGAAGGTCTACGGCAAAAAGGATGGAGTATCCTCAATCTACTGGTTATTCAAAGAGAGGCTAAAATCAG AAACACTGCTGTCAAAGTCTTGGCACAAGGTGATAATCAAGTTATTTGCACACAGTATAAAACGAAGAAATC ${\tt GAGAAACGTTGTAGAATTACAGGGTGCTCTCAATCAAATGGTTTCTAATAATGAGAAAATTATGACTGCAAT}$ CAAAATAGGGACAGGGAAGTTAGGACTTTTGATAAATGACGATGAGACTATGCAATCTGCAGATTACTTGAA TTATGGAAAAATACCGATTTTCCGTGGAGTGATTAGAGGGTTAGAGACCAAGAGATGGTCACGAGTGACTTG TGTCACCAATGACCAAATACCCACTTGTGCTAATATAATGAGCTCAGTTTCCACAAATGCTCTCACCGTAGC TCATTTTGCTGAGAACCCAATCAATGCCATGATACAGTACAATTATTTTGGGACATTTGCTAGACTCTTGTT GATGATGCATGATCCTGCTCTTCGTCAATCATTGTATGAAGTTCAAGATAAGATACCGGGCTTGCACAGTTC TACTTTCAAATACGCCATGTTGTATTTGGACCCTTCCATTGGAGGAGTGTCGGGCATGTCTTTGTCCAGGTT ${\tt TTTGATTAGAGCCTTCCCAGATCCCGTAACAGAAAGTCTCTCATTCTGGAGATTCATCCATGTACATGCTCG}$ AAGTGAGCATCTGAAGGAGATGAGTGCAGTATTTGGAAACCCCGAGATAGCCAAGTTTCGAATAACTCACAT AGACAAGCTAGTAGAAGATCCAACCTCTCTGAACATCGCTATGGGAATGAGTCCAGCGAACTTGTTAAAGAC TGAGGTTAAAAAATGCTTAATCGAATCAAGACAAACCATCAGGAACCAGGTGATTAAGGATGCAACCATATA

Figure 14 continued

 ${\tt TTTGTATCATGAAGAGGATCGGCTCAGAAGTTTCTTATGGTCAATAAATCCTCTGTTCCCTAGATTTTTAAG}$ TGAATTCAAATCAGGCACTTTTTTGGGAGTCGCAGACGGGCTCATCAGTCTATTTCAAAATTCTCGTACTAT TCGGAACTCCTTTAAGAAAAAGTATCATAGGGAATTGGATGATTTGATTGTGAGGAGTGAGGTATCCTCTTT GACACATTTAGGGAAACTTCATTTGAGAAGGGGATCATGTAAAATGTGGACATGTTCAGCTACTCATGCTGA TCCACAACATCGAAAAGAGACTCCTTGTGCACCATGTAACACATCAGGGTTCAATTATGTTTCTGTGCATTG TCCAGACGGGATCCATGACGTCTTTAGTTCACGGGGACCATTGCCTGCTTATCTAGGGTCTAAAACATCTGA ATCTACATCTATTTTGCAGCCTTGGGAAAGGGAAAGCAAAGTCCCACTGATTAAAAAGAGCTACACGTCTTAG AGATGCTATCTCTTGGTTTGTTGAACCCGACTCTAAACTAGCAATGACTATACTTTCTAACATCCACTCTTT ${\tt ATCTCGGATGAGCCATGGTGGGTTCGCATCTCAGAGCACTGCAGCATTGACCAGGTTGATGGCAaCTACAGA}$ $\tt CACCATGAGGGATCTGGGAGATTTCGACTTTTATTCCAGGCAACGTTGCTCTATGCTCAGATTAC$ CACCACTGTTGCAAGAGACGGATGGATCACCAGTTGTACAGATCATTATCATATTGCCTGTAAGTCCTGTTT GAGACCCATAGAAGAGATCACCCTGGACTCAAGTATGGACTACACGCCCCCAGATGTATCCCATGTGCTGAA GACATGGAGGAATGGGGAAGGTTCGTGGGGACAAGAGATAAAACAGATCTATCCTTTAGAAGGGAATTGGAA ${\tt GAATTTAGCACCTGCTGAGCAATCCTATCAAGTCGGCAGATGTATAGGTTTTCTATATGGAGACTTGGCGTA}$ $\tt CTCCTATCCGACAAGCAACCGTGATATGGGGGTGATTGTCAGAAATTACTTCAAATACCAATGCCGTCTAAT$ TGAAAAGGGAAAATACAGATCACATTATTCACAATTATGGTTATTCTCAGATGTCTTATCCATAGACTTCAT TGGACCATTCTCTATTTCCACCACCCTCTTGCAAATCCTATACAAGCCATTTTTATCTGGGAAAGATAAGAA ATTCTTCACCAAGGACATATTATTGTGTCCAGAGGAAATCAGACATGCTTGCAAGTTCGGGATTGCTAAGGA TAATAATAAAGACATGAGCTATCCCCCTTGGGGAAGGGAATCCAGAGGGACAATTACAACAATCCCTGTTTA TTATACGACCACCCTTACCCAAAGATGCTAGAGATGCCTCCAAGAATCCAAAAATCCCCTGCTGTCCGGAAT CAGGTTGGGCCAGTTACCAACTGGCGCTCATTATAAAATTCGGAGTATATTACATGGAATGGGAATCCaTTA CAGGGACTTCTTGAGTTGTGGAGACGGCTCCGGAGGGATGACTGCTGCATTACTACGAGAAAATGTGCATAG TGCCCTAGAAACTTTAGGAGGAGATAAATCGAGATGTGTAAATGGTGAAACATGTTGGGAATATCCATCTGA $\tt CTTATGTGACCCAAGGACTTGGGACTATTTCCTCCGACTCAAAGCAGGCTTGGGGCTTCAAATTGATTTAAT$ TGTAATGGATATGGAAGTTCGGGATTCTTCTACTAGCCTGAAAATTGAGACGAATGTTAGAAATTATGTGCA $\tt CCGGATTTTGGATGAGCAAGGGTTTTAATCTACAAGACTTATGGAACATATATTTGTGAGAGCGAAAAGAA$ TGCAGTAACAATCCTTGGTCCCATGTTCAAGACGGTCGACTTAGTTCAAACAGAATTTAGTAGTTCTCAAAC GTCTGAAGTATATATGGTATGTAAAGGTTTGAAGAAATTAATCGATGAACCCAATCCCGATTGGTCTTCCAT

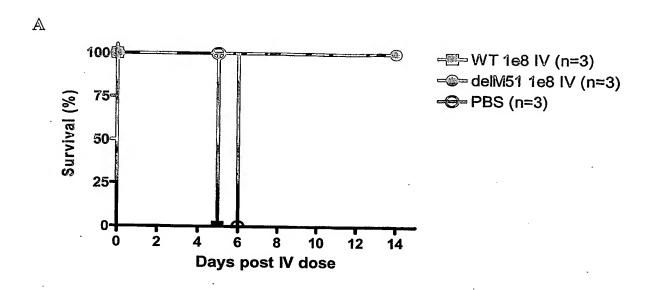
Figure 14 continued

Figure 14 continued

Nucleic Acid Sequence of the M Protein Gene for VSV Mutant AV2

Amino Acid Sequence for the M Protein of VSV Mutant AV2

MSSLKKILGLKGKGKKSKKLGIAPPPYEEDTNMEYAPSAPIDKSYFGVDEMDTHDPHQLRYEKFFFTVKMTV RSNRPFRTYSDVAAAVSHWDHMYIGMAGKRPFYKILAFLGSSNLKATPAVLADQGQPEYHAHCEGRAYLPHR MGKTPPMLNVPEHFRRPFNIGLYKGTVELTMTIYDDESLEAAPMIWDHFNSSKFSDFREKALMFGLIVEKKA SGAWFLDSVRHFK



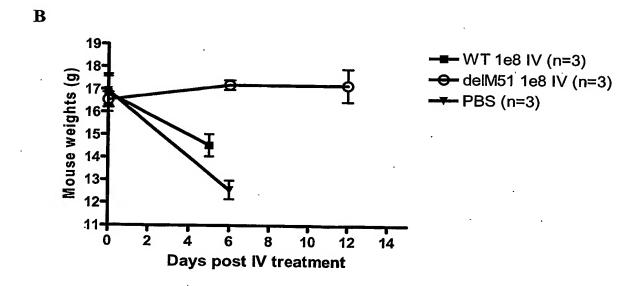


FIGURE 17A-B

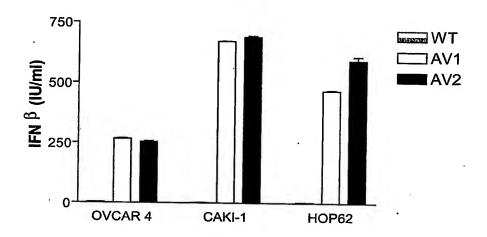
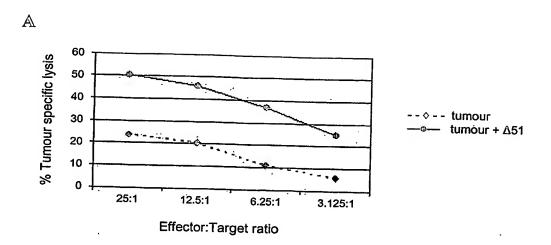


FIGURE 18



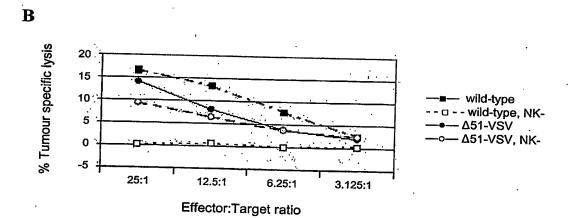


FIGURE 19

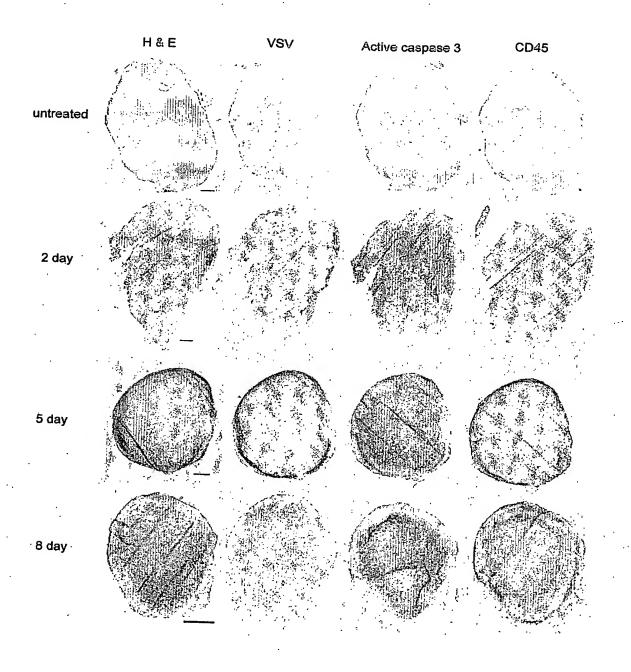


FIGURE 20